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Reviewer: Anne Corrigan

Timestamp: Thu Sep 13 18:13:05 EDT 2007

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Application No: 10578234 Version No: 1.0

Input Set:

Output Set:

Started: 2007-09-04 13:36:46.672

Finished: 2007-09-04 13:36:47.304

Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 632 ms

Total Warnings: 0

Total Errors: 0

No. of SeqIDs Defined: 24

Actual SeqID Count: 24

SEQUENCE LISTING

<110> CuraGen Corporation et al.
 Alsobrook II, John P.
 Eichen, Joseph
 Lepley, Denise M.
 Miller, Charles E.
 Mezes, Peter

<120> COMPOSITIONS AND METHODS OF USE FOR A FIBROBLAST GROWTH FACTOR

<130> 15966-557 SNP NATL (Cura-57 SNP NATL)

<140> 10578234

<141> 2007-09-04

<150> 10/702,126

<151> 2003-11-04

<160> 24

<170> CuraSeqList version 0.1

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Ile Leu Glu Phe Ile Ser Val Ala Val Gly Leu Val Ser Ile Arg Gly	
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Val Asp Ser Gly Leu Tyr Leu Gly Met Asn Asp Lys Gly Glu Leu Tyr	
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gga tca gag aaa ctt act tcc gaa tgc atc ttt agg gag cag ttt gaa	432
Gly Ser Glu Lys Leu Thr Ser Glu Cys Ile Phe Arg Glu Gln Phe Glu	
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gag aac tgg tat aac acc tat tca tct aac ata tat aaa cat gga gac	480
Glu Asn Trp Tyr Asn Thr Tyr Ser Ser Asn Ile Tyr Lys His Gly Asp	
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Thr Gly Arg Arg Tyr Phe Val Ala Leu Asn Lys Asp Gly Thr Pro Arg	
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gat ggc gcc agg tcc aag agg cat cag aaa ttt aca cat ttc tta cct	576
Asp Gly Ala Arg Ser Lys Arg His Gln Lys Phe Thr His Phe Leu Pro	
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aga cca gtg gat cca gaa aga gtt cca gaa ttg tac aag gac cta ctg	624
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Val Asp Ser Gly Leu Tyr Leu Gly Met Asn Asp Lys Gly Glu Leu Tyr
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 Thr Tyr Ser Ser Asn Ile Tyr Lys His Gly Asp Thr Gly Arg Arg Tyr
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Gln Gly Thr Arg Gln Asp His Ser Leu Phe Gly Ile Leu Glu Phe Ile
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Tyr Leu Gly Met Asn Asp Lys Gly Glu Leu Tyr Gly Ser Glu Lys Leu
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Thr Ser Glu Cys Ile Phe Arg Glu Gln Phe Glu Glu Asn Trp Tyr Asn
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Thr Tyr Ser Ser Asn Ile Tyr Lys His Gly Asp Thr Gly Arg Arg Tyr
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Phe Val Ala Leu Asn Lys Asp Gly Thr Pro Arg Asp Gly Ala Arg Ser
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Leu Gly Gln Pro Gly Ala Ala Gln Leu Ala His Leu His Gly Ile Leu	
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cgc cgc cgg cag ctc tat tgc cgc acc ggc ttc cac ctg cag atc ctg	144
Arg Arg Arg Gln Leu Tyr Cys Arg Thr Gly Phe His Leu Gln Ile Leu	
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ccc gac ggc agc gcg cag ggc acc cgg cag gac cac agc ctc ttc ggt	192
Pro Asp Gly Ser Ala Gln Gly Thr Arg Gln Asp His Ser Leu Phe Gly	
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Ile Leu Glu Phe Ile Ser Val Ala Val Gly Leu Val Ser Ile Arg Gly	
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Gly Ser Glu Lys Leu Thr Ser Glu Cys Ile Phe Arg Glu Gln Phe Glu	
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gag aac tgg tat aac acc tat tca tct aac ata tat aaa cat gga gac	384
Glu Asn Trp Tyr Asn Thr Tyr Ser Ser Asn Ile Tyr Lys His Gly Asp	
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act ggc cgc agg tat ttt gtg gca ctt aac aaa gac gga act cca aga	432
Thr Gly Arg Arg Tyr Phe Val Ala Leu Asn Lys Asp Gly Thr Pro Arg	
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Asp Gly Ala Arg Ser Lys Arg His Gln Lys Phe Thr His Phe Leu Pro	
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aga cca gtg gat cca gaa aga gtt cca gaa ttg tac aag aac cta ctg	528
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Pro Asp Gly Ser Ala Gln Gly Thr Arg Gln Asp His Ser Leu Phe Gly
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Ile Leu Glu Phe Ile Ser Val Ala Val Gly Leu Val Ser Ile Arg Gly
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Val Asp Ser Gly Leu Tyr Leu Gly Met Asn Asp Lys Gly Glu Leu Tyr
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Gly Ser Glu Lys Leu Thr Ser Glu Cys Ile Phe Arg Glu Gln Phe Glu
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Glu Asn Trp Tyr Asn Thr Tyr Ser Ser Asn Ile Tyr Lys His Gly Asp
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Thr Gly Arg Arg Tyr Phe Val Ala Leu Asn Lys Asp Gly Thr Pro Arg
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